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Title: MD Simulations of Full-Scale HIV-1 Lipid Envelope on LANL Grizzly Supercomputer

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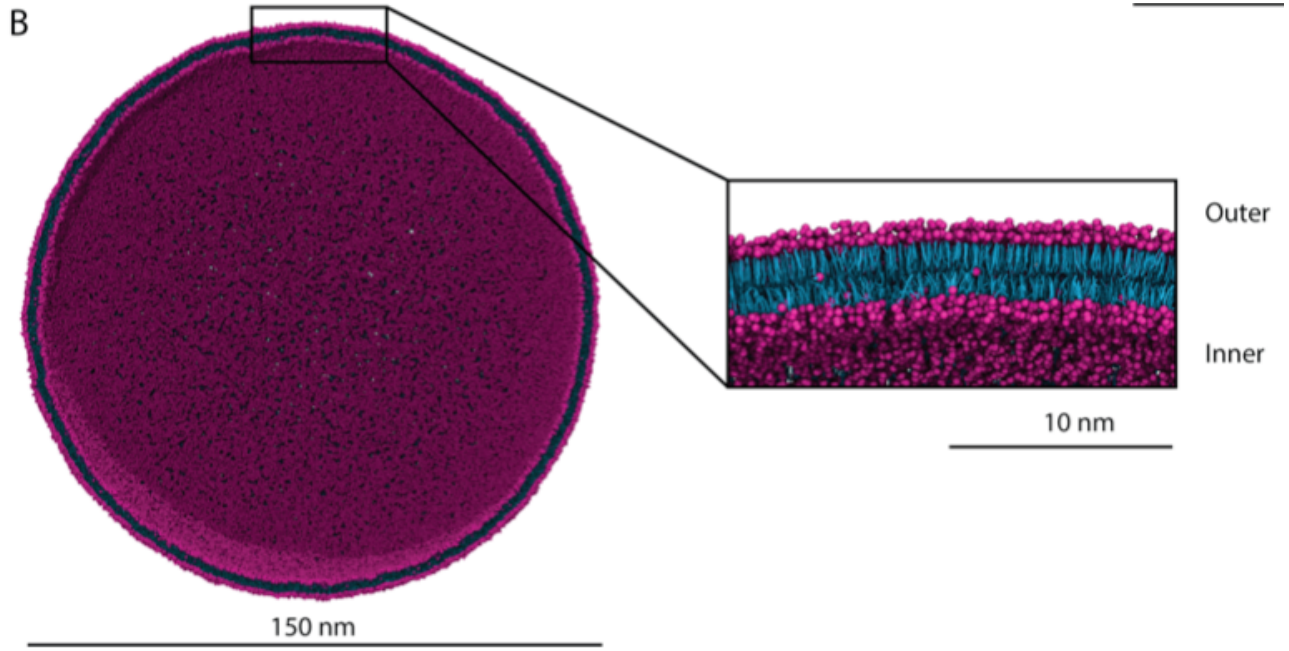
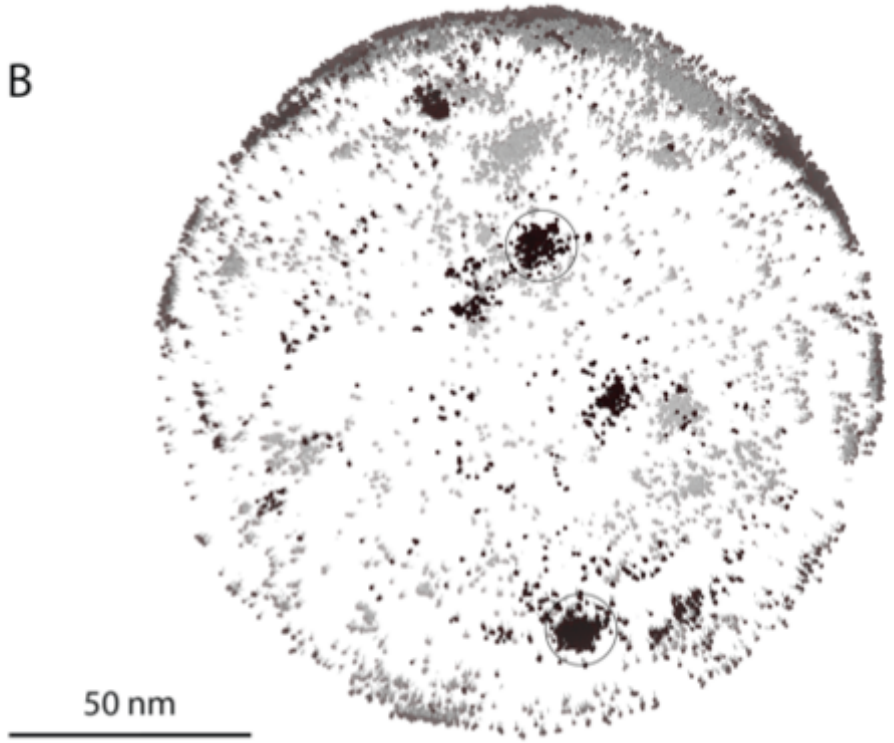
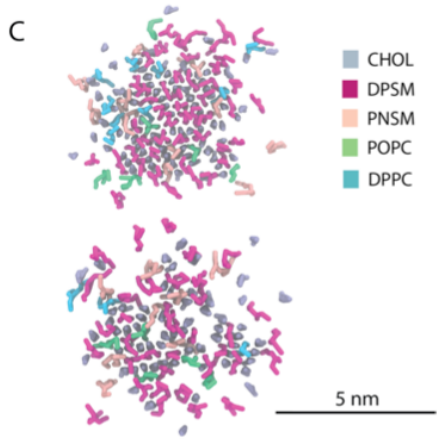


Cross-section view of the HIV-1 vesicle after 5 microseconds of simulation with the MARTINI force field.



Determination of low-mobility regions in HIV-1 vesicle (circled) over the course of the simulation.



Identification of Lipid Species
In Low Mobility Regions

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